SEQUENCE LISTING

	Singh, Mohan Bhalla, Prem Hui-Ling, Xu Swoboda, Ines	
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tct gat ata aat tgc tcc Ser Asp Ile Asn Cys Ser 125		agttggca gagaataatc 4	198
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Ile	Ile	Arg	Pro	Leu 85	Ser	Asp	Gly	Thr	Asn 90	Asn	Cys	Leu	Val	Asn 95	Asn	
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Lys Gln Pro Thr Thr Thr Ser Gly Lys Trp Arg Phe Ala Arg Phe 50 55 60

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- 27 -

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: (OTHER THAN US): THE UNIVERSITY OF MELBOURNE (US ONLY): SINGH Mohan, BHALLA Prem, HUI-LING Xu and

SWOBODA Ines

(ii) TITLE OF INVENTION:

NOVEL NUCLEIC ACID MOLECULES AND USES

THEREFOR

(iii) NUMBER OF SEQUENCES: 9

(iv) CORRESPONDENCE ADDRESS:

- (A) ADDRESSEE: DAVIES COLLISON CAVE
- (B) STREET: 1 LITTLE COLLINS STREET
- (C) CITY: MELBOURNE
- (D) STATE: VICTORIA
- (E) COUNTRY: AUSTRALIA
- (F) ZIP: 3000

(v) COMPUTER READABLE FOR M:

- (A) MEDIUM TYPE: Floppy dick
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATIÓN DATA:

- (A) APPLICATION NUMBER: PCT INTERNATIONAL
- (B) FILING DATE: 24-JUL-1998
- (C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

- (A) APPLICATION NUMBER: PO8233
- (B) FILING DA/TE: 25-JUL-1997
- (C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

- (A) APPLICATION NUMBER: PP1184
- (B) FILING DATE: 31-DEC-1997
- (C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

- (A) NAME: HUGHES, DR E JOHN L
- (C) REFERENCE/DOCKET NUMBER: EJH/AF

- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: +61 3 9254 2777
 - (B) TELEFAX: +61 3 9254 2770
 - (C) TELEX: AA 31787

															/	
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	(i)	(A) (B) (C)	ENCE C LENGT TYPE: STRAN TOPOL	H: 1 ami DEDN	4 am no a ESS:	ino cid sin	acid	S				/				
	(ii)	MOLE	CULE T	YPE:	pep	tide					,					
	(xi)	SEQU.	ENCE D	ESCR:	IPTI(ON:	SEQ :	ID NO	0:1:							
GTA	CTCTI	'AA GC	ATACAA	CA TO	GAG						,					14
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	(i)	(A) (B) (C)	ENCE CH LENGTH TYPE: STRANI TOPOLO	H: 14 amir DEDNE	ami no ac ESS:	ino a cid sing	acids	5	/							
	(ii)	MOLEC	CULE TY	PE:	pept	ide	/	/					1,00			
													પ્રાપ્યો કેવે ઉદ્ય			
	(xi)	SEQUE	NCE DE	ESCRI	PTIC	N: 5	SKO 1	D NC):2:						11.24	
CAGG	CATA	CT TGA	ATGCT <i>I</i>	AC AF	\GA		,						ť.		1.	14
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	(ix)		IRE: NAME/I LOCAT			. 468										
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GGGT	FACTC	AA TT	CATAC	AA C	ATG Met	AGG Arg	GCG Ala	GTG Val	GCG Ala 5	GTT Val	TTC Phe	TTT Phe	GCT Ala	TGC Cys 10		111
GTT Val	CTC Leu	TTC/TC Phe Cy	GT ATG ys Met 15	GTT Val	CAC His	AAA Lys	GCC Ala	GCA Ala 20	CTT Leu	GCG Ala	GAT Asp	GAT Asp	AAA Lys 25	ACG Thr		159
TGC	VVC	CET AC	CA GAT	ттт	ATG	GTT	ACC	CAA	ACC	ATA	ACT	GGA	TTG	ACA		207

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TTAA	TGGA	AT G	AATTE	TGTA	T T	AGAAT	TGA	AAA A	(AAT)	AAA	AAA	LAAA.	AAA A	AAA	λΑΑΑ		615
AAAA	AAAA	AA									•						625
121	TNEC	NDM X II	T ON	EOD	CEO	TD A	10.4.										
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	(3	ii) N	OLEC	CULE	TYPI	E: pi	rote	in									
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578 587

(2) INFORMATION FOR SEQ ID NO:6:

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	(i	ii) 1	MOLEC	ULE	TYPE	: pr	otei	ln								
	()	(i) :	SEQUE	NCE	DESC	RIPT	'ION:	SEC	DID	NO: 6	5:					
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Leu	Arg	Ser	Glu 20	Lys	Ala	λla	Leu	Gln 25	Phe	Ser	Val	Ser	Arg 30	Val	Glu	
Tyr	Ser	Leu 35	Lys	Lys	Gly	Arg	Tyr 40	Cys	Arg	Arg	Leu	Gly 45	Ala	Thr	Ala	
Pro	Val 50	Туг	Leu	Ala	Ala	Val 55	Leu	Glu	Asn	Leu	Val 60	Ala	Glu	Val	Leu	
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Pro	Arg	His	Ile	Met 85	Leu	Ala	Pro	Arg	Asn 90	Asp	Val	Glu	/Val	Glu 95	Gln	
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(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 111 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: protein	
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Ala Ser Lys Ala Leu Arg Lys Ala Pro Pro Pro Pro Thr Lys Gly Val	
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His Arg Lys Leu Pro Phe Gln Gly Leu Wal Arg Lys Ile Trp Gln Asp 65 70 75 80	
Leu Lys Thr His Leu Arg Phe Lys Asm His Ser Val Pro Pro Leu Glu 85 90 95	
Glu Val Thr Glu Val Tyr Pro Cys Gln Thr Ile Gly Gly Cys Tyr 100 105 110	

1	2) INFORMATION	FOR	SEO	TD	NO . 9 .

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 945 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA

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(2) INFORMATION FOR SEQ ID NO:10:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH 14 base pairs

CAGTGGTGTT TCTTTTTGCT TCCGTTCTCT TCTGTATGGT TCACA

- (B) TYPE: hucleic acid
- (C) STRANDEDNESS: single (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA
- (xi) SEQUENCE/DESCRIPTION: SEQ ID NO:10:

CAGGCATACT TGAATCCTAC AAGA

- (2) INFORMATION FOR SEQ ID NO:11:
 - (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs

14

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- 35 -

- (B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

TGTGAACCAT ACAGAAGAGA ACGC